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_ <del>z</del>√. ∌
                          DGENE
     ACA50276 DNA
     New antisense nucleic acids, useful for identifying proteins or screening
ΤI
      for homologous nucleic acids required for cellular proliferation to
      isolate candidate molecules for rational drug discovery programs -
      Wang L; Zamudio C; Malone C; Haselbeck R; Ohlsen K L; Zyskind J W; Wall
IN
      D; Trawick J D; Carr G J; Yamamoto R; Forsyth R A; Xu H H
                  ELITRA PHARM INC.
PA
PΙ
      WO 2002077183
                      A2 20021003
                                                 999
      WO 2002-US9107
ΑI
                           20020321
      US 2001-815242
                           20010321
PRAI
      US 2001-948993
                           20010906
      US 2001-342923P
                           20011025
      US 2002-72851
                           20020208
      US 2002-362699P
                           20020306
PSL
      Claim 14; SEQ ID No 38146
DED
      19 JUN 2003 (first entry)
DT
      Patent
LA
      English
      2003-029926 [02]
OS
      P-PSDB: ABU46406
CR
DESC Prokaryotic essential gene #31933.
      Antisense; ds; prokaryotic essential gene; cell proliferation; drug
KW
      design; gene.
ORGN
     Streptococcus pyogenes.
      The invention relates to an isolated nucleic acid comprising any one of
AΒ
      the 6213 antisense sequences given in the specification where expression
      of the nucleic acid inhibits proliferation of a cell. Also included are:
      (1) a vector comprising a promoter operably linked to the nucleic acid
      encoding a polypeptide whose expression is inhibited by the antisense
      nucleic acid; (2) a host cell containing the vector; (3) an isolated
      polypeptide or its fragment whose expression is inhibited by the
      antisense nucleic acid; (4) an antibody capable of specifically binding
      the polypeptide; (5) producing the polypeptide; (6) inhibiting cellular
      proliferation or the activity of a gene in an operon required for
      proliferation; (7) identifying a compound that influences the activity of
      the gene product or that has an activity against a biological pathway
      required for proliferation, or that inhibits cellular proliferation; (8)
      identifying a gene required for cellular proliferation or the biological
      pathway in which a proliferation-required gene or its gene product lies
      or a gene on which the test compound that inhibits proliferation of an
      organism acts; (9) manufacturing an antibiotic; (10) profiling a
      compound's activity; (11) a culture comprising strains in which the gene
      product is overexpressed or underexpressed; (12) determining the extent
      to which each of the strains is present in a culture or collection of
      strains; or (13) identifying the target of a compound that inhibits the
      proliferation of an organism. The antisense nucleic acids are useful for
      identifying proteins or screening for homologous nucleic acids required
      for cellular proliferation to isolate candidate molecules for rational
      drug discovery programs, or for screening homologous nucleic acids
      required for proliferation in cells other than S. aureus, S. typhimurium,
      K. pneumoniae or P. aeruginosa. The present sequence is one of the target
      prokaryotic essential genes. Note: The sequence data for this patent did
      not form part of the printed specification, but was obtained in
      electronic format directly from WIPO at ftp.wipo.int/pub/published pct_se
      quences.
NΑ
      188 A; 101 C; 109 G; 139 T; 0 other
SQL
      537
SEQ
        1 atqtcacqta ttqqtaataa agtaattact atgcctgcag gcgttgaatt
       51 aacaaataac aacaatgtta ttactgttaa aggccctaaa ggcgaactca
      101 ctcgtgagtt caacaaaaat attgaaatca aagttgaagg gactgaaatc
      151 acagttgtac gtcctaacga ctcaaaagaa atgaaaacaa tccatggtac
      201 aacccqtqct aacttqaata acatqqttqt aqqtqtttct qaaqqtttca
      251 aaaaagatet tgaaatgaag ggtgteggtt accgtgetea actteaaggt
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301 actaaacttg tcctttcagt aggtaaatct caccaagacg aagttgaagc 351 tccagaagga attactttca ctgttgctaa cccaacttca atctcagttg 401 aaggaatcaa caaagaagtt gttggtcaaa cagctgctta catccgtagc 451 ttgcgttcac cagagcctta caaaggcaaa gggatccgtt acgttggtga 501 atacgtacgc cttaaagaag gtaaaacagg taaataa